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Attorney Docket No. ORT-1414

IN THE UNITED STATES PATENT & TRADEMARK OFFICE

-----X-----

Re: U.S. PATENT APPLICATION
Serial Number 09/833,222

Filed: April 11, 2001

Art Unit 1647

-----X-----

DECLARATION OF DR. NING QI UNDER RULE 37 CFR 1.131

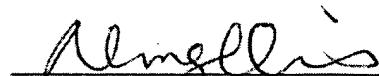
1. I am a co-inventor of the application Serial Number 09/833,222, entitled "cDNA Encoding the Human Alpha2Delta-4 Calcium Channel Subunit" (the '222 application). I obtained my Ph.D. degree in July 1993, from Baylor College of Medicine, and had my postdoctoral training at University of California, Los Angeles from 1994-1997. From 1997 to 1999, I was employed as a research assistant professor at UCLA and then, as an Assistant Professor at University of Kentucky. I have been employed by Johnson & Johnson as a research scientist since September 1999. My resume is attached as Exhibit 1.
2. The invention described in the '222 application is based on the research my colleague and I conducted from 1999 to 2000, during which we successfully cloned the human $\alpha_2\delta$ -4 calcium channel subunit, as evidenced by the data in my laboratory notebook. I attach the relevant notebook pages (Exhibits 2 and 3) and a computer printout (Exhibit 4) herewith to demonstrate the process of our invention. This was completed long before the Brown and Bertelli's publication (March 22, 2001, WO 01/19870A2).

3. I was interested in cloning this subunit because the voltage gated calcium channel is one of important pain therapeutic targets as recognized by the scientists in the field of pain control. I believe that the $\alpha_2\delta$ is a key regulatory subunit and that there are multiple $\alpha_2\delta$ isoforms, based on our analysis on human genomic DNA data, all play an important role in mediating Ca^{2+} influx in excitable cells.
4. I started the project of cloning $\alpha_2\delta-4$ on or about September 20, 1999 (Exhibit 12. No. 15798-1) and my colleague and I continuously worked on the project for the next several months. Because the cDNA encoding $\alpha_2\delta-4$ subunit is more than 3Kb, I had to clone it piece by piece. (Exhibit 2). We ultimately assembled and sequenced the fully length cDNA of $\alpha_2\delta-4$ subunit by April 7, 2000, which was recorded in my lab notebook (Exhibit 3, p. No. 15798-139), and the DNA and protein sequence information were also documented in both Microsoft word and Seqweb files on the same day. (Exhibit 4).

I declare under the penalty of perjury that the above statements are true and accurate to the best of my knowledge.

Dated: June 6, 2006

By:


Ning Qin, Ph.D.

CURRICULUM VITAE

[04-30-2006]

Ning Qin

SITE: Spring House

DIVISION: Analgesics

DEPARTMENT: Drug Discovery

EDUCATION: Baylor College of Medicine, Houston, Texas
1993, Biochemistry, Ph.D.

EMPLOYMENT

1999-present Johnson & Johnson Pharmaceutical Research and Development
Spring House, PA
2003-Present Principal Scientist, Analgesics, Drug Discovery
1999-2002 Senior Scientist

1999- Department of Pharmacology, University of Kentucky
Lexington, KY 40536
1999-1999 Assistant Professor

1994-1998 Dept. of Anesthesiology UCLA. School of Medicine,
Los Angeles, CA 90095
1997-1998 Research Assistant Professor
1994-1996 Postdoctoral Fellow

CURRENT PROFESSIONAL AFFILIATIONS

1. Society of Neuroscience
2. Biophysical Society
3. American Pain Society
4. International Association for the study of Pain
5. New York Academy of Sciences

PUBLICATIONS

Manuscripts

1. N. Qin, S. J. Pittler and W. Baehr, (1992) "In Vitro Isoprenylation and Membrane Association of Mouse Photoreceptor cGMP Phosphodiesterase α and β subunits Expressed in Bacteria", *Journal of Biological Chemistry*, 267, 8458-63
2. N. Qin and W. Baehr, (1993) "Expression of Mouse Rod Photoreceptor PDE γ Subunit in Bacteria", *FEBS Letters*, 321, 6-10
3. M. L. Suber, S. J. Pittler, N. Qin, G. C. Wright, V. Holcombe, R. H. Lee, C. M. Craft, R. N. Lolley, W. Baehr and R. L. Hurwitz, (1993) "Irish Setter Dogs Affected with Rod/Cone Dysplasia Contain a Nonsense Mutation in the Rod cGMP Phosphodiesterase β Subunit Gene", *Proc. Natl. Acad. Sci. U.S.A.* 90, 3968-3972
4. N. Qin and W. Baehr, (1994) "Expression and Mutagenesis of Biologically Active Mouse Rod Photoreceptor cGMP Phosphodiesterase", *Journal of Biological Chemistry*, 269 3265-3271
5. T. Schneider, X. Wei, R. Olcese, J. L. Costantin, A. Neely, P. Palade, E. Peryes, N. Qin, J. Zhou, G. Crawford, R. G. Smith, S. H. Appel, E. Stefani and L. Birnbaumer (1994) "Molecular analysis and functional expression of the human type E neuronal Ca^{2+} -channel α_1 subunit", *Receptors and Channels*, 2, 255-270
6. R. Olcese, N. Qin (*Co-first author*), T. Schneider, A. Neely, X. Wei, E. Stefani and L. Birnbaumer (1994) "The Amino Terminus of a Calcium Channel β subunit Sets Rates of Channel Inactivation Independently of the Subunit's Effect on Activation", *Neuron*, 13, 1433-1438
7. F. Noceti, P. Baldelli, X. Wei, N. Qin, L. Toro, L. Birnbaumer and E. Stefani (1996) "Effective Gating Charges Per Channel in Voltage Dependent K^+ and Ca^{2+} Channels", *J. Gen. Physiol.* 108, 143-155
8. N. Qin, R. Olcese, J. Zhou O. Caballo, L. Birnbaumer and E. Stefani (1996) "Identification of a second region of the β subunit involved in regulation of calcium channel inactivation", *Am. J. Physiol. (Cell Physiology)*, 271:C1539-C1545
9. R. Olcese, A. Neely, N. Qin, X. Wei, L. Birnbaumer and E. Stefani (1997) "Coupling Between Charge Movement and Pore Opening in Neuronal α_{1E} Calcium Channel." *J. Physiol.* 497, 675-686

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11. J. Zhou R. Olcese, N. Qin, E. Stefani and L. Birnbaumer, (1997) "Feedback inhibition of calcium channels by calcium depends on a short sequence of the carboxyterminus that does not include the Ca^{2+} binding function of a motif with similarity to calcium binding domains", *Proc. Natl. Acad. Sci. U.S.A.* 94, 2301-2305
12. N. Qin, D. Platano, R. Olcese, E. Stefani and L. Birnbaumer (1997) "Inhibition of Neuronal Ca^{2+} Channel G-protein Coupled receptors Mediated by a Small C-terminal $\text{G}\beta\gamma$ Binding Domain of α_1 ", *Proc. Natl. Acad. Sci. U.S.A.* 94, 8866-887.
13. J. Costantin, F. Noceti, N. Qin, X. Wei, L. Birnbaumer and E. Stefani, (1998) "Facilitation by the β Subunit of Pore Openings in Cardiac Ca^{2+} Channels", *J. Physiol.* 507, 93-103
14. F. Noceti, R. Olcese, N. Qin, and E. Stefani, (1998) " Ca^{2+} -dependent inactivation in $\alpha_{1\text{C}}$ Ca^{2+} channels: effect of Bay K 8644(-) and the $\beta_{2\text{a}}$ subunit. A model", *J. Gen. Physiol.* 111, 463-475
15. N. Qin, R. Olcese, E. Stefani and L. Birnbaumer (1998) "Modulation of human neuronal $\alpha_{1\text{E}}$ type calcium channel by $\alpha_2\delta$ subunit", *Am. J. Physiol. (Cell Physiology)*, 274, C1324-C1331
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17. N. Li, R.N. Fariss, K. Zhang, A. Otto-Bruc, F. Haeseleer, D. Bronson, N. Qin, A. Yamazaki, I. Subbaraya, A. H. Milam, K. Palczewski and W. Baehr, (1998) "Guanylate-cyclase-inhibitory protein is a frog retina Ca^{2+} binding protein related to mammalian guanylate-cyclase-activating protein", *Eur. J. Biochem.* 252, 591-59
18. J. Costantin, N. Qin, L. Birnbaumer and E. Stefani, (1998) "Long lasting voltage-dependent facilitation of a cardiac and brain calcium channel: correlation with the coupling efficiency between charge movement and pore opening", *FEBS Letter*, 423, 213-217
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20. J. Costantin, N. Qin, M. N. Waxham, L. Birnbaumer and E. Stefani, (1999) "Complete reversal of run-down in rabbit cardiac Ca^{2+} channels by patch-cramming in *Xenopus* oocytes; partial reversal by protein kinase A". *Pflugers Arch*; 437(6): 888-94
21. N. Qin, R. Olcese, M. Bransby, T. Lin, and L. Birnbaumer (1999) " Ca^{2+} -induced inhibition of the cardiac Ca^{2+} channel depends on calmodulin". *Proc. Natl. Acad. Sci. U.S.A.*; 96(5): 2435-8
22. B. Vannier, M. Peyton, G. Boulay, D. Brown, N. Qin, M. Jiang, X. Zhu, and L. Birnbaumer (1999) "Mouse trp2, the homologue of the human trpc2 pseudogene, encodes mTrp2, a store depletion-activated capacitative Ca^{2+} entry channel" *Proc. Natl. Acad. Sci. U.S.A.*; 96(5): 2060-4

23. G. Boulay, D. M. Brown, N. Qin (*Co-first author*), M. Jiang, A. Dietrich, M. X. Zhu, Z. Chen, M. Birnbaumer, K. Mikoshiba, and L. Birnbaumer (1999) "Modulation of Ca^{2+} entry by polypeptides of the inositol 1,4,5-trisphosphate receptor (IP3R) that bind transient receptor potential (TRP): Evidence for roles of TRP and IP3R in store depletion-activated Ca^{2+} entry" *Proc. Natl. Acad. Sci. U S A*; 96: 14955-14960.
24. D. Platano, N. Qin, F. Noceti, L. Birnbaumer, E. Stefani, and R. Olcese (2000) "Expression of alpha2delta Subunit Interferes with Prepulse Facilitation in Cardiac L-type Calcium Channels" *Biophys. J.* 78 2959-2972
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26. N. Qin, S. Yagel, M. Momplaisir, E. E. Codd and M. R. D'Andrea (2002) "Molecular cloning and characterization of human voltage-gated calcium channel $\alpha_2\delta$ -4 subunit" *Molecular Pharmacology*, 62 (3)
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28. N. Qin, M. D'Andrea, M. Lubin, N. Shafae, E. Codd and A. Correa (2003) "Cloning and Characterization of a novel splicing variant of human sodium channel β_1 subunit". *The FEBS Journal (European Journal of Biochemistry)*, 270: 4762-4770
29. R. Pagani, M. Song, M. McEnery, N. Qin, R. W. Tsien, L. Toro, E. Stefani and O. D. Uchitel (2004) "Differential expression of alpha 1 and beta subunits of voltage dependent Ca^{2+} channel at the neuromuscular junction of normal and P/Q Ca^{2+} channel knockout mouse". *Neuroscience*: 123 (1): 75-85
30. N. Qin, SP Zhang, T.L. Reitz, J.M. Mei and C.M. Flores (2005) "Cloning, Expression and Functional Characterization of Human COX-1 Splicing Variants: Evidence for Intron 1 Retention" *The Journal of Pharmacology and Experimental Therapeutics*, 315 (3): 1298-1305
31. Y. Liu, M. Lubin, T.L. Reitz, Y. Wang, R.W. Colburn, C.M. Flores and N. Qin (2006) "Molecular identification and functional characterization of a temperature-sensitive transient receptor potential channel (TRPM8) from canine" *European Journal of Pharmacology*, 530: 23-32
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Book Chapters

1. L. Birnbaumer, N. Qin, R. Olcese, E. Tareilus and E. Stefani, (1998). "Studies on the regulation of the human neuronal calcium channel by β and $\alpha_2\delta$ subunits". *Low voltage activated T-type calcium channels*, 258-268, Montpellier, France

2. F. Noceti, R. Olcese, P. Baldelli, N. Qin, L. Birnbaumer and E. Stefani, (1998) "Charges per channel in Ca^{2+} channels: effect of the regulatory β subunit co-expression on α_1 pore forming subunit". *Low voltage activated T-type calcium channels*, 307-313, Montpellier, France
3. R. Olcese, F. Noceti, P. Baldelli, N. Qin, L. Birnbaumer and E. Stefani, (1998) " β subunit modulation of the coupling between charge movement and pore opening in calcium channels". *From Ion Channels to Cell to Cell Conversation*, 91-104, Plenum press, New York and London
4. L. Birnbaumer, N. Qin, R. Olcese, E. Tareilus, Daniela Platano and E. Stefani, (1998) Structures and Functions of Calcium Channel β Subunits." *J. of Bioenergetics and Biomembranes*, Vol. 30(4) 357-375
5. L. Birnbaumer, G. Boulay, D. Brown, M. Jiang, A. Doerlich, K. Mikoshiba, X. Zhu, and N. Qin, (2000) "Mechanism of Capacitative Ca^{2+} entry (CCE): Interaction between IP3 Receptor and TRP links the Internal Calcium Storage Compartment to Plasma Membrane CCE Channels." *Recent Progress in Hormone Research*, Vol. 55: 127-162

ABSTRACTS

1. Y. Liu, M.L. Lubin, T. Reitz, R.W. Colburn, C.M. Flores, and N. Qin "Cloning and functional characterization of a canine transient receptor potential channel, cTRPM8". 2004 Neuroscience Meeting
2. N. Qin, S-P. Zhang, T. Reitz, E. E. Codd and C. M. Flores "Human COX-3? Cloning and Characterization of a splicing variant of COX-1" 2004 Neuroscience meeting
3. N. Qin, M. Nepper, Y. Liu and C. M. Flores "Characterization of calcium channel beta subunit interaction domain" 2005 International Association for the Study of Pain Annual Meeting

PATENTS

1. N. Qin, E. Codd, and M. D'Andrea (2006) "The Human Voltage Gated Sodium Channel β_{1A} Subunit And Method Of Use." **US 6994,993 B2**
2. N. Qin and E. Codd (2003) "cDNA encoding the human alpha2delta-4 calcium channel subunit" **US20030166045 A1**
3. N. Qin, E. Codd C. M. Flores and SP. Zhang (2004) "Human cyclooxygenase-3 enzyme and uses thereof" **US20040235017 A1**
4. C. M. Flores, Y. Liu, M. Lubin and N. Qin (2006) "Canine Cold- And Menthol-Sensitive Receptor 1" **US20060014246 A1**
5. N. Qin, Y. Liu, M. Nepper, T. Reitz and C. M. Flores (2005) "Compositions and Methods for Identifying Modulators of TRPV2" Provisional filing
6. N. Qin, Y. Liu and C. M. Flores (2005) "A Polypeptide Complex of TRPM8 and Calmodulin And Its Uses Thereof" Provisional filing

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R.W. Johnson Pharmaceutical Research Institute

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Department Anesthetics Code 62333

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Project No. *α2S of cat channel* Protocol / Experiment No. Date *9/21/99*
 Subject Purpose *cloning*

Contd. from page _____

*Search Genebank for all published Calcium channels
 916 documents*

NCBI Entrez Nucleotide QUERY BLAST Entrez ?

Current Query

Details Search : calcium channel[All Fields] --> Retrieve 916 Documents
 Number of documents to display per page: Mod. Date limit: No Limit

Add Term(s) to Query :

Search Field: All Fields Search Mode: Automatic Search Clear

Enter Terms: _____

Enter one or more author last names, text words, or other keywords. To search for all terms that begin with a given word, place an asterisk (*) at the end of the word. Journal Titles must be MEDLINE abbreviations; Author names must be in the form LastName Initial(s), e.g. Smith BJ. The initials can be omitted. Detailed Help is available.

 Clear All

Modify Current Query :

Term (Total Records)

calcium channel[All Fields] (916)
 Search for the Intersection (AND) of the selected terms.

Questions or comments? Write to the NCBI Help Desk.

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Investigator *Wing Oin*Date *9/28/99*Read and Understood *Ellen Odd*Date *10/7/99*

No. 15798- 6

Project No. α₂S Protocol / Experiment No. Date 9/28/99

Subject Purpose

Contd. from page 5

Analysis and Comparison of other small fragments of α₂S identified on 9/20/99.

Two fragments are novel α₂S

name them to α₂S-4.

they are H86016

g1436938.

Two are human α₂S-3 fragments.

they are g3805345

g2885043. or AA001473

sequence comparison results of H86016
& g1436938

are saved in

H:/mydata/sene/α₂S/new-α₂S.

Contd. on page 5

Investigator Dr. S

Date 10/1/99

Read and Understood E. Codd

Date 10/07/99

No. 15798- 8

Project No. α_2S Protocol / Experiment No. Date 7/30/89

Subject Purpose

Contd. from page 7

1. comparison of 208672.2
199408.3

W:66 mouse α_2S -3.

same as H:/mydata/gene/1a2d/Inayte clone 208672
Inayte clone 199408.3

2. Design oligos.

α_2-3-5

-6
-7
-8
-9
-10

} for clone human α_2S -3.

α_2-4-1

-2
-3
-4
-5

} for clone human α_2S -4.

Contd. on page 6

Investigator mg

Date 10/1/89

Read and Understood E Gold

Date 10/07/99

Project No. 15798-2 No. 31A Protocol / Experiment No. Date 10-2-99

Subject Purpose

Contd. from page

1. Race PCR to amplify NT of h2S-4.
 Marathon Ready Library ① fetal human brain
 ② human brain.

RX. 5ul 10x Buffer.

5ul DNA library.

1ul AP1.

1ul A2-4-9 (10 pmol/ul).

1ul Helvorne Tag.

1ul 10mM dNTP.

35ul H₂O

cycles. 94°C / 30''

↓
94°C / 5''] 5 cycles.↓
72°C / 4 min↓
94°C / 5''] 5 cycles.↓
70°C / 4 min↓
94°C / 5''] 25 cycles.↓
68°C / 4 min↓
4°C / parking.Contd. on page 22Investigator DRWDate 11/16/99Read and Understood DRWDate 3-6-00

Project No. 15798-28 Protocol / Experiment No. Date 11-8-99

Subject Purpose

Contd. from page _____

1. cloning Race PCR products (11-5-99) of NT α -8-4 (human)(1) gel purification
cut two parts.

A. 3 kb band

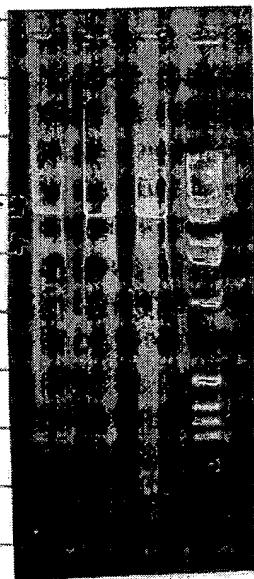
B. 2-2.5 kb bands

purify them
separately

(2) patch them.

(3) Ligatin' script.

(4) transfer to XL10-gold

2. Retest condition for Na- β A.

RX 50 ml. w/o Salvage Tag

(1) SB1-4 + SB1-2. } rat brain cDNA

(2) SB1-5 + SB1-2 }

(3) SB1-1 + SB1-2 } human brain cDNA

(4) SB1-5 + SB1-2 }

condition: 30 cycles.

 $94^{\circ}\text{C}/30'' \rightarrow [58^{\circ}\text{C}/30''] \rightarrow 72^{\circ}\text{C}/60''$

5 ml to check PCR RX.

does not work.

Contd. on page 75Investigator DR J WDate 11/16/99Read and Understood W/WDate 3-6-00

Project No. 128 Protocol / Experiment No. Date 11-11-99
 Subject Purpose

Contd. from page _____

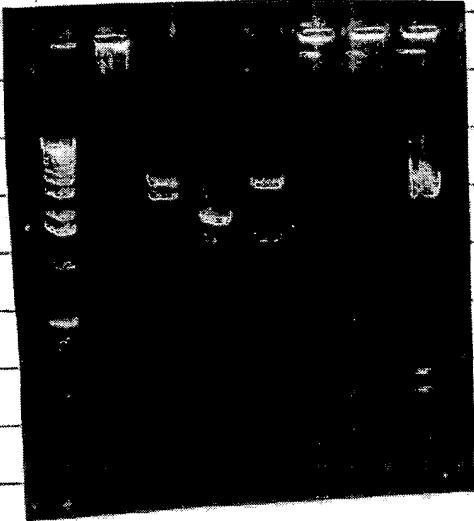
1. miniprep NQC44.
 two new 3 Kb α 28-4 in PCR 2.1
 four new 2 kb α 28-4 in PCR 2.1
 dig. NQC44 with EcoRI + KpnI
 all others with EcoRI only
 $3^1 3^2 2^1 2^2 2^3 2^4$, NQC44.

3-2, 2-1, & 2-2

have insertion

3-2 \rightarrow NQC452-1 \rightarrow NQC462-2 \rightarrow NQC47

Send NQC44-47

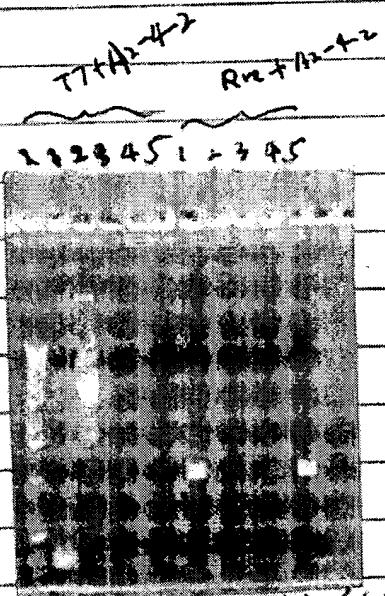
for seq.

2. characterize these clones. by PCR.

- ① NQC44, ② NQC45, ③ NQC46
- ④ NQC47, ⑤ (maybe 3343341)

20 μ l RX.NQC44 & 45 are very promising
 but 44 is too short.

45 is right size.

Contd. on page 32Investigator DJWRead and Understood DNWDate 11/16/99Date 3-6-00

Project No. 928 Protocol / Experiment No. Date 11-11-99
 Subject Purpose

Contd. from page _____

1. miniprep NQC44.
 two new 3 kb α S-4 in PCR 2.1
 four new 2 kb α S-4 in PCR 2.1
dig. NQC44 with EcoRI + KpnI
all others with EcoRI only
3' 1 3' 2 2' 2 2' 3' 4' NQC44.

3-2, 2-1, 2 2-2

have insertion

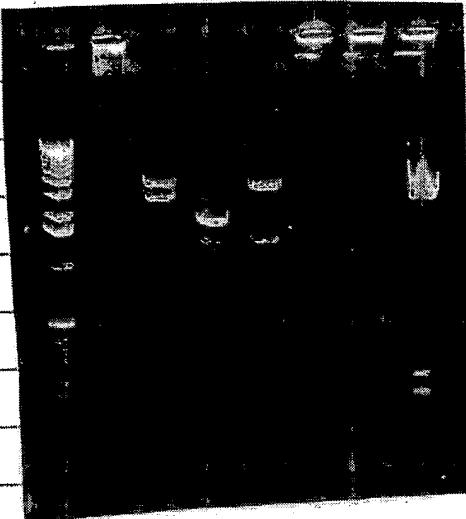
3-2 \rightarrow NQC45

2-1 \rightarrow NQC46

2-2 \rightarrow NQC47

Send NQC44-47

for seq.



2. characterize these clones. by PCR.

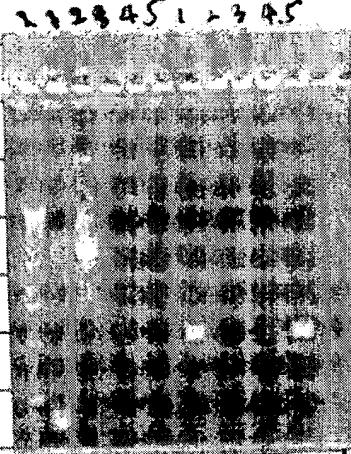
- ① NQC44, ② NQC45, ③ NQC46
- ④ NQC47, ⑤ maybe 3343341.

20 μ l RX.

NQC44 & 45 are very promising
 but 44 is too short.

45 is right size.

$T1 + A + H + T$
 $R + H + T + H$



Contd. on page 32

Investigator DR

Date 11/16/99

Read and Understood W/W

Date 3-6-00

Project No. DN-β / Cu-αβ Protocol / Experiment No. Date 4/11/00
Subject Purpose

Contd. from page _____

2. Analysis of ~~soy~~ Data of h₂S-4 NT.

They (WAC 100-195) are all h₂S-4 NT.

first shell clones of h₂S-4.

Syn Ab. No.: A2-4-27 } for add HA on h₂S-4 CT
A2-4-28 }

A2-3-29 } add HA on
A2-3-30 } h₂S-3 CT.

Contd. on page 137Investigator DR Date 4/14/00Read and Understood DRDate 4/11/00

ha2d-4b.pep

[Text View](#)

!!AA_SEQUENCE 1.0

ha2d-4.pep Length: 1090 April 7, 2000 14:19 Type: P Check: 9269 ..

1 MAVALGTRRR DRVKLWADTF GGDLYNTVTK YSGSLLLQKK YKDVESSLKI
51 EEVGDGLELVR KFSEDMENML RRKVEAVQNL VEAEEAADLN HEFNESLVFD
101 YYNSVLLINER DEKGNFVELG AEFLLESNAH FSNLPVNTSI SSVQLPTNVY
151 NKDPDILNGV YMSEALNAVF VENFQRDPTL TWQYFGSATG FFRIYPGIKW
201 TPDENGVITF DCRNRGWIYQ AATSPKDIVI LVDVSGSMKG LRMTIAKHTI
251 TTILDTLGEN DFVNIIAYND YVHYIEPCFK GILVQADRDN REHFKLLVEE
301 LMVKGVGVVD QALREAFQIL KQFQEAKQGS LCNQAIMLIS DGAVEDYEPV
351 FEKYNWPDCK VRVFTYLIGR EVSFADRMKW IACNNKGYYT QISTLADTQE
401 NVMEYLHVLS RPMVINHDHD IIWTEAYMDS KLLSSQAQSL TLLTTVAMPV
451 FSKKNETRSH GILLGVVGSD VALRELMKLA PRYKLGVHGY AFLNTNNGYI
501 LSHPDLRPLY REGKKLKPKP NYNSVLDSEV EWEDQAESLR TAMINRETGT
551 LSMDVKVPMDF KGKRVLFLLTN DYFFTDISDT PFSLGAVLSR GHGEYILLGN
601 TSVEEGLHDL LHPDLALAGD WIYCITDIDP DHRKLSQLEA MIRFLTRKDP
651 DLECDEELVR EVLFDAVVTA PMEAYWTALA LNMSEESEHV VDMAFLGTRA
701 GLLRSSLFVG SEKVSDRKFL TPEDEASVFT LDRFPLWYRQ ASEHPAGSFV
751 FNLRWAEGPE SAGEPMVVT A STAVAVTV DK RTAIAAAAGV QMKLEFLQRK
801 FWAATRQCST VDGPYTQSCE DSDLDCFVID NNGFILISKR SRETGRFLGE
851 VDGAVLTQLL SMGVFSQV TM YDYQAMCKPS SHHHSAAQPL VSPISAF LT A
901 TRWLLQELVL FLLEWSVWGS WYDRGAEAKS VFHHSHKHKK QDPLQPCDTE
951 YPVFVYQPAI REANGIVECG PCQKVFVVQQ IPNSNLLLV TDPTCDCSIF
1001 PPVLQEATEV KYNASVKCDR MRSQKLRRRP DSCHAFHPEV RVEADRGWAG
1051 FSSPNPLCLG LCPCRQEHI G MPMNTPVPVL LGGNIRVYAL

Human $\alpha_2\delta 4$ map

April 7, 2000 14:04

Map of a DNA sequence showing restriction enzyme cleavage sites and a splice site. The sequence is numbered 1 to 451. The map includes labels for HindIII, BtsI, NcoI, PstI, BspGI, DrdII, and EcoRI, along with their corresponding restriction sites. A splice site is marked with "Splicing A2-4-37, HindIII". The sequence is numbered 1 to 451, and the map includes a scale bar from 0 to 540.

Key features include:

- HindIII:** Cuts at positions 1, 90, and 180.
- BtsI:** Cuts at positions 1, 91, and 181.
- NcoI:** Cuts at positions 1, 91, and 181. A splice site is marked at position 181.
- PstI:** Cuts at positions 1, 91, 181, 271, and 361.
- BspGI:** Cuts at positions 1, 91, 181, 271, 361, and 451.
- DrdII:** Cuts at positions 1, 91, 181, 271, 361, and 451.
- EcoRI:** Cuts at positions 1, 91, 181, 271, 361, and 451.
- Splicing A2-4-37, HindIII:** Located between positions 181 and 271.
- Scale:** 0 to 540.

NarT

DRAFT

A2-4-24
A2-4-29
541 CTCGGGCCGAGTTCCCTGGAGTCCAATGCTCactTTCAAGCAACCTGCGGtGAAACACCTtCATCAGGAGCTGGCAGGCTGGCCCAACC
631 CTCGACCCGGGCTCAAGGGAGGACCTCAAGGTtTACGAGGtGAAGTGGTGGACGGCCactTTGCGACGTTGACGGGGTGG
E L G A E F I L E S N A H F S N L P V N T S I S S V Q L P T -
BsrgI |
DraI |
PciI |
BsmI |
AACGTGTACACAAAGACCCAGATTTAAATGGAGTCTACATGCTGAAAGCCTTGAATGCTGTCCTGGAGAGAACTTCCAGAGAGAC
631 TTGACATGTTGTTCTGGGTCTATAAAATTACCTCAAGACTTCAAATGATGTCAGACCTACAGACAGAACCTTGAAGGTCTCTCTG
N V Y N K D P D I L N G V Y M S E A L N A V F V E N F Q R D -

721 CCAACGTTGACCTGGCAATATTTGGCAGTGCACCTGGATTCTAGGATCTATCCAGGTATAAAATGGACACCTGatTGAGAATGGAGTC
 P T L T W Q Y F G S A T G F F R I Y P G I K W T P D E N G V -
 BtsI SspI BtsI
 A2-4-19

	BamHI	DraI	BspGI	BspGI
991	TACAAATGACTACGTCCATTACATCGAGCCTTGTAAAGGGATCCTGTCaGGGGACCGAGACATCGAGAGCATTCAAATGCTGCTG			
	ATGTTACTGTAGTCAGGTAAATGTTAGCTGGTAAACAAATTCCCTAGGAGCAGGTCCGGCTGGCTCTGTAGCTCTCGTAAGGTTGACGAC			
	Y N D Y V H Y I E P C F K G I V Q A D R D N R E H F K L L -			
	A2-4-32			
1081	GTGAGGAGTGTGATGGTCAAAGGTGTGGGGGTCTGAGAAGGCCCTGAGAAGCCTTCAGATCCAGTCCAGTCAGCAGTCCAGAGGCCAAAG			
	CACTCTCAACTACCAGTTCCACACCCCCAGCACCTGGACTCTCTGGAAAGGTCTAGGACTTCGTCAGGTTCTCGGTTCAGGTTCTCGGTTC			
	V E E L M V K G V G V D O A L R E A F O I I K O F A K -			

Narrator: Pfiff 111087
MSCI

Map of a DNA sequence with restriction enzyme cleavage sites and restriction fragments labeled A2-4-1 through A2-4-30. The map includes labels for enzymes SphI, NcoI, Bpu11I, PmlI, BspGI, StuI, BtsI, and HindIII, and markers Q, G, S, L, C, N, O, A, I, M, L, I, S, D, G, A, V, E, P, V, F, T, Y, L, I, G, R, E, V, S, F, A, D, R, M, K, W, I, A, C, N, N, K, G.

Key features of the map:

- Enzymes and cleavage sites:**
 - SphI:** A2-4-11, A2-4-12, A2-4-13.
 - NcoI:** A2-4-14, A2-4-15, A2-4-16, A2-4-17, A2-4-18, A2-4-19.
 - Bpu11I:** A2-4-14.
 - PmlI:** A2-4-14.
 - BspGI:** A2-4-29.
 - StuI:** A2-4-18.
 - BtsI:** A2-4-17.
 - HindIII:** A2-4-30.
- Restriction fragments:** A2-4-1 through A2-4-30.
- Marker positions:** Q, G, S, L, C, N, O, A, I, M, L, I, S, D, G, A, V, E, P, V, F, T, Y, L, I, G, R, E, V, S, F, A, D, R, M, K, W, I, A, C, N, N, K, G.

BsrGI

1711 CCCCTGTACAGAGGGGAGAACTAAACCAAACCTAACTACAACAGTGTGGATCTCCGAAGGGAGTGGGAAGACAGGGCTGAA
 GGGACATGTCCTCCCTTCTTGTGATGGATGTCACACCTAGAGGGCTTACCTCACCCCTACCTCGTCCGACTT
 P L Y R E G K K P K N Y N S V D L S E V E W D Q A E -
 TCTCTGAGAACAGCCATGATCAATAGGGAAACAGGTACTCTCGATGGATGTGAAGGTTCCGATGGATAAAGGGAAAGCGAGTTCTTTC
 AGAGACTCTTGTGGTACTAGTTATCCCTTGTCCATGAGAGGCTACACTTCCAAGGGTACCTATTCCCTTCGCTCAAGAAAAG
 S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F -

Sma I

1891 CTGACCAATGACTACTTTACGGACATCAGGACACCCCTTCACTGGGGTGGTCTGACTTGGCCCTGGGAAATACATCCCT
 GACTGGTTACTGTGAAAGGAAGTGCCTGTAGTGCCTGAGGGAAAGTCAAACCCCAACCGACAGGGCCGGTGCCTCTATGTAGGAA
 L T N D Y F F T D I S D T P F S L G V V L S R G H G E Y I L -

BsrI

1981 CTGGGAACACGCTCTGTGAAAGGAAGGCTGCACTGACTTGGCCCTGGCCCTGGCTGACTGGATCTACTGGATCACAGAT
 GACCCCTTGTGAGACACCTCTCCGGACGTACTGAAACGAAAGTGGTCTGGAACCGGGACCGGCCACTGACCTAGATGACGTAGTGTCTA
 L G N T S V E E G L H D L L H P D L A G D W I Y C I T D -

2071 ATTGACCCAGAACCCACGGCTCAGCCAGCTAGAGGCCATGATCCGGCTTCCCTACCCAGGAAGGACCCAGACCTGGAGTGTGACGGAG
 TAACTGGGTCTGGGCCCTTCGAGTCGGTCACTCCGGTACTAGGGGAAGGGTGGTCTGGACCTCACACTGCTCCCT
 I D P D H R K L S Q L E A M I R F L T R K D P D L E C D E E -

Nco I

2161 CTGGTCCGGAGGTGCTGTTGACCCGGTGGTGACAGCCCCATGGAAAGCCATGGCCTACTGGCACGGCTGGCTGGCCCTCAACATGTCGGAGGACT
 GACCAAGGCCCTCCACGACAAACTGGCCACCACTGTGGGGTACCTTCGGATGACCTGTGCGAACGGGAGTTGACAGGGCTCTCAGA
 L V R E V L F D A V V T A P M E A Y W T A L A L N M S E E S -

Pml I

2251 GAACACGTGGTGGACATGGCCTCCCTGGCACCCGGCTGGCCCTCTGAGAACGGCTTGTGGCTGGGCTCCGAGAACGGTCTCCGACAGG
 CTTGTGCACCACTGGAAAGGACCCGTGGGCCGACCGGAGGACTCTTCGTGCGAACAAAGCACCGAGGGCTCTCCAGAGGCTGTCC
 E H V V D M A F L G T R A G L L R S S L F V G S E D R -

Sma I

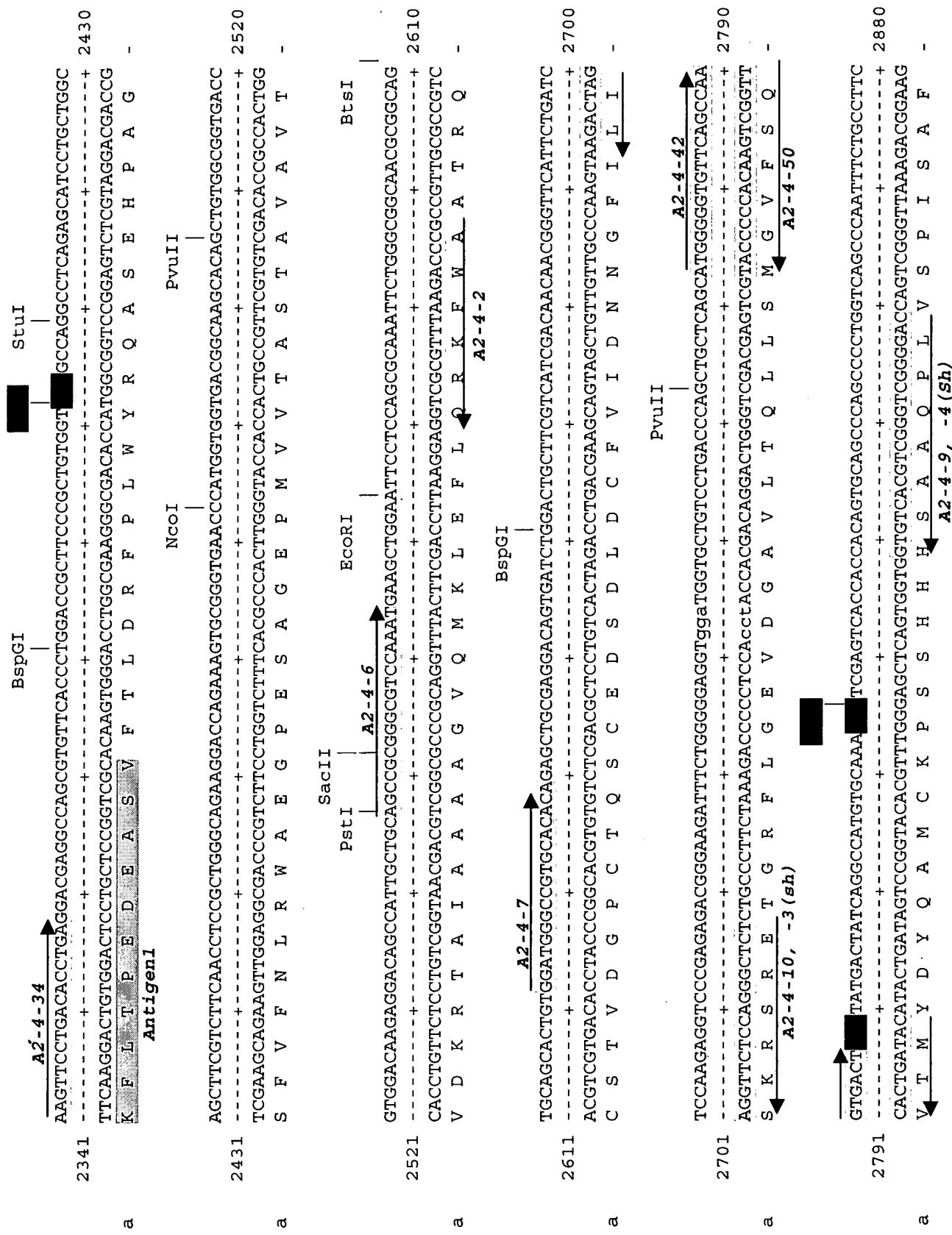
1800 CCCCTGTACAGAGGGGAGAACTAAACCAAACCTAACTACAACAGTGTGGATCTCCGAAGGGAGTGGGAAGACAGGGCTGAA
 GGGACATGTCCTCCCTTCTTGTGATGGATGTCACACCTAGAGGGCTTACCTCACCCCTACCTCGTCCGACTT
 P L Y R E G K K P K N Y N S V D L S E V E W D Q A E -
 TCTCTGAGAACAGCCATGATCAATAGGGAAACAGGTACTCTCGATGGATGTGAAGGTTCCGATGGATAAAGGGAAAGCGAGTTCTTTC
 AGAGACTCTTGTGGTACTAGTTATCCCTTGTCCATGAGAGGCTACACTTCCAAGGGTACCTATTCCCTTCGCTCAAGAAAAG
 S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F -

Sma I

1980 CCCCTGTACAGAGGGGAGAACTAAACCAAACCTAACTACAACAGTGTGGATCTCCGAAGGGAGTGGGAAGACAGGGCTGAA
 GGGACATGTCCTCCCTTCTTGTGATGGATGTCACACCTAGAGGGCTTACCTCACCCCTACCTCGTCCGACTT
 P L Y R E G K K P K N Y N S V D L S E V E W D Q A E -
 TCTCTGAGAACAGCCATGATCAATAGGGAAACAGGTACTCTCGATGGATGTGAAGGTTCCGATGGATAAAGGGAAAGCGAGTTCTTTC
 AGAGACTCTTGTGGTACTAGTTATCCCTTGTCCATGAGAGGCTACCTATTCCCTTCGCTCAAGAAAAG
 S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F -

Sma I

2340 CCCCTGTACAGAGGGGAGAACTAAACCAAACCTAACTACAACAGTGTGGATCTCCGAAGGGAGTGGGAAGACAGGGCTGAA
 GGGACATGTCCTCCCTTCTTGTGATGGATGTCACACCTAGAGGGCTTACCTCACCCCTACCTCGTCCGACTT
 P L Y R E G K K P K N Y N S V D L S E V E W D Q A E -
 TCTCTGAGAACAGCCATGATCAATAGGGAAACAGGTACTCTCGATGGATGTGAAGGTTCCGATGGATAAAGGGAAAGCGAGTTCTTTC
 AGAGACTCTTGTGGTACTAGTTATCCCTTGTCCATGAGAGGCTACCTATTCCCTTCGCTCAAGAAAAG
 S L R T A M I N R E T G T L S M D V K V P M D K G K R V L F -



T-5

DNA Sequence:

1 CAGGTACATT CAGCAGAGCC CAAGTCTGCC ACTCTCCAAC CaGAGGCCCT
51 GGAAGCTTGG GGTCAAGCTC AGTCCTGGC TCGTCAGCCC GGCCCCACAA
101 CCCTCAGCAG GAGaACCTGC CGAGGACATT CAGCACACAG CAGTGCAGCC
151 GCTGGGTCTCT GAGGGTTCTC CGCGTCTCCT GCCCAGGCCA TGGCTGTAGC
201 TTTAGGGACA AGGAGGAGGG ACAGAGTGAA GCTATGGGCT GACACCTTCG
251 GCGGGGACCT GTATAACAct GTGACCAAAT ACTCAGGCTC TCTCTTGCTG
301 CAgAAGAAGT ACAAGGATGT GGAGTCCAGT CTGAAGATCG AGGAGGTGGA
351 TGGCTTGGAG CTGGTGAGGA AGTTCTCAGA GGACATGGAG AACATGCTGC
401 GGAGGAAAGT CgAGGCGGTC CAgAATCTGG TGGAAGCTGC CGAGGAGGCC
451 GACCTGAACC ACGAATTCAA TGAATCCCTG GTGTTCGACT ATTACAACCTC
501 GGTCTTGATC AACGaGAGGG ACGAGAAGGG CaACTTcGTG GAGCTGGCG
551 CCGAGTCCCT CCTGGAGTCC AATGCTCaCT TCAGCAACCT GCCGGtGAAC
601 ACCTcCATCA GCAGCGTGCA GCTGCCACC AACGTGTACA ACAAAGACCC
651 AGATATTTA AATGGAGTCT ACATGTCTGA AgCCTTGAAT GCTGTCTTCG
701 TGGAGAACTT CCAGAGAGAC CCAACGTTGA CCTGGCAATA TTTTGGCAGT
751 GCAACTGGAT TCTTCAGGAt CTATCCAGGT ATAAAATGGA CACCTGATGA
801 GAATGGAGTC ATTACTTTG ACTGCCGAAA CCGCGGCTGG TACATTCAAG
851 CTGCTACTTC TCCCAAGGAC ATAGTGATTT TGGTGGACGT GAGCGGCAGT
901 ATGAAGGGGC TGAGGATGAC TATTGCCAG CACaCCATCA CCACCATCTT
951 GGACACCCTG GGGGAGAATG ACTTCGTTAA TATCATAGCG TACAATGACT
1001 ACGTCCATTA CATCGAGCCT TGTTTAAAG GGATCCTCGT CCAGGCGGAC
1051 CGAGACAATC GAGAGCATT CAAGACTgCTG GTGGAGGAGT TGATGGTCAA
1101 AgGTGTGGGG GTCGTGGACC AAGCCCTGAG AGAAGCCTC CAGATCCTGA
1151 AgCAGTTCCA AGAgGCCAAG CAAGGAAGCC TCTGCAACCA GGCATCATG
1201 CTCATCAGCG ACgGCGCCGT GGAGGACTAC GAGCCGGTGT TTGAGAAGTA
1251 TAACTGGCCA GACTGTAAGG TCCGAGTTT CACTTACCTC ATTGGGAGAG
1301 AAGTGTCTTT TGCTGACCGC ATGAAGTGGG TTGCATGCAA CAACAAAGGC
1351 tACTACACGC AGATCTCAAC GCTGGCGGAC ACCCAGGAGA ACGTGATGGA
1401 ATACCTGCAC GTGCTCAGCC GCCCCATGGT CATCAACCAC GACCACGACA
1451 TCATCTGGAC AGAGGCCCTAC ATGGACAGCA AGCTCCTCAG CTCGCAGGCT
1501 CAGAGCCTGA CACTGCTCAC CACTGTGGCC ATGCCAGTCT TCAGCAAGAA
1551 GAACGAAACG CGATCCCCTG GCATTCTCCT GGGTGTGGTG GGCTCAGATG
1601 TGGCCCTGAG AGAGCTGATG AAGCTGGCGC CCCGGTACAA GCTTGGAGTG
1651 CACGGATAACG CCTTTCTGAA CACCAACAAT GGCTACATCC TCTCCCATCC
1701 CGACCTCCGG CCCCTGTACA GAGAGGGGAA GAAACTAAAA CCCAAACCTA
1751 ACTACAAACAG TGTGGATCTC TCCGAAGTGG AGTGGGAAGA CCAGGCTGAA
1801 TCTCTGAGAA CAGCCATGAT CAATAGGGAA ACAGGTAATC TCTCGATGGA
1851 TGTGAAGGTT CCGATGGATA AAGGGAGCG AGTTCTTTT CTGACCAATG
1901 ACTACTTCTT CACGGACATC AGGGACACCC CTTTCAGTTT GGGGGCGGTG
1951 CTGTCCCAGG GCCACGGAGA ATACATCCTT CTGGGAAACA CGTCTGTGGA
2001 AGAAGGCCTG CATGACTTGC TTCACCCAGA CCTGGCCCTG GCCGGTGA
2051 GGATCTACTG CATCACAGAT ATTGACCCAG ACCACCGGAA GCTCAGCCAG
2101 CTAGAGGCCA TGATCCGCTT CCTCACCAAGG AAGGACCCAG ACCTGGAGTG
2151 TGACGAGGAG CTGGTCCGGG AGGTGCTGTT TGACGCGGTG GTGACAGCCC
2201 CCATGGAAGC CTACTGGACA GCGCTGGCCC TCAACATGTC CGAGGAGTCT
2251 GAACACGTGG TGGACATGGC CTTCTGGC ACCCGGGCTG GCCTCCTGAG
2301 AAGCAGCTTG TTCGTGGGCT CCGAGAAGGT CTCCGACAGG AAGTTCTGAA
2351 CACCTGAGGA CGAGGCCAGC GTGTTCACCC TGGACCGCTT CCCGCTGTGG
2401 TACCGCCAGG CCTCAGAGCA TCCCTGCTGGC AGCTTCGTCT TCAACCTCCG
2451 CTGGGCAGAA GGACCAGAAA GTGCGGGTGA ACCCATGGTG GTGACGGCAA
2501 GCACAGCTGT GGCGGTGACC GTGGACAAGA GGACAGCCAT TGCTGCAGCC
2551 GCGGGCGTCC AAATGAAGCT GGAATTCCCTC CAGCGCAAAT TCTGGGCGGC
2601 AACGGCCAG TGCAGCACTG TGGATGGGCC GTACACACAG AGCTGCGAGG
2651 ACAGTGATCT GGACTGCTTC GTCATCGACA ACAACGGTT CATTCTGATC
2701 TCCAAGAGGT CCCGAGAGAC GGGAAAGATT CTGGGGGAGG TggaTGTTGC
2751 TGTCTTGACC CAGCTGCTCA GCATGGGGGT GTTCAGCCAA GTGACTATGT
2801 ATGACTATCA GGCCATGTGC AAACCCCTCGA GTCACCCACCA CAGTGCAGCC
2851 CAGCCCCCTGG TCAGCCCCAT TTCTGCCTTC TTGACGGCGA CCAGGTGGCT
2901 GCTGCAGGAG CTGGTGTGTT TCCTGCTGGA GTGGAGTGTG TGAGGCTCCT
2951 GGTACGACAG AGGGGGcgaG GCCAAAAGTG TCTTCCATCA CTCCCCACAAA
3001 CACAagaagC AGGACCCGCT GCagcCCTGc gacaCGgagt ACCccgTGTt
3051 cGTGTAccaG CGGGccatTCC GGGaggCCAA CGGGATCGTG GAGTGCAGGGC

3101 CCTGCCAGAA GGTATTTGTG GTGCAGCAGA TTCCCAACAG TAACCTCCTC
 3151 CTCCCTGGTGA CAGACCCAC CTGTGACTGC AGCATCTTCC CACCAGTGCT
 3201 GCAGGAGGCG ACAGAAGTCA AATATAATGC CTCTGTCAAA TGTGACCGGA
 3251 TGCCTCTCCA GaagctccGC CGGCGACCAG ACTCCTGCCA CGCCTTCCAT
 3301 CCAGAGGTGC GGGTTGAGGC GGATCGAGGG TGGGCTGGAT TTTCATCCCC
 3351 AAACCCCTCTG TGCCTGGTC TGTGCCCCTG CAGACAGGAG CATATAGGGA
 3401 TGCCAATGAA CACACCTGTG CCTGTGCTTC TCGGGGGAAA CATTGCGT
 3451 TATGCCCTGT GACACTGTGA TATAATAAGA AACAGA

Protein Sequence:

V	T	K	M	A	V	A	L	G	T	R	R	R	D	R	V	K	L	W	A	D	T	F	G	D	L	Y	N	T	
L	V	R	K	F	S	E	D	M	E	N	M	L	R	R	K	V	E	S	S	L	K	I	E	V	D	G	L	E	E
D	L	N	H	E	F	N	E	S	L	V	F	D	Y	Y	N	S	V	L	I	N	E	R	D	E	K	G	N	F	
E	L	G	A	E	F	L	L	E	S	N	A	H	F	S	N	L	P	V	N	T	S	I	S	S	V	E	N	F	
N	V	Y	N	K	D	P	D	I	L	N	G	V	Y	M	S	E	A	L	N	A	V	F	V	E	N	F	R	D	
P	T	L	T	W	Q	Y	F	G	S	A	T	G	F	F	R	I	Y	P	G	I	K	W	T	P	D	E	N	G	
I	T	F	D	C	R	N	R	G	W	Y	I	Q	A	F	A	T	S	P	K	D	I	V	I	L	V	D	V	S	
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D	C	K	V	R	V	F	T	Y	L	I	G	R	E	V	S	F	A	D	R	M	K	W	I	A	C	N	N	K	
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D	H	D	I	I	W	T	E	A	Y	M	D	S	K	L	L	S	S	Q	A	Q	S	L	T	L	T	T	V	A	
M	P	V	F	S	K	K	N	E	T	R	S	H	G	I	L	L	G	V	V	G	S	D	T	V	A	L	R	E	
K	L	A	P	R	Y	K	L	G	V	H	G	Y	A	F	L	N	T	N	N	G	Y	I	L	S	H	P	D	L	R
P	L	Y	R	E	G	K	K	L	K	P	K	P	N	Y	N	S	V	D	L	S	E	V	W	E	D	Q	A	E	
S	L	R	T	A	M	I	N	R	E	T	G	T	L	S	M	D	V	K	V	P	M	D	K	G	K	R	V	L	
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I	D	P	D	H	R	K	L	S	Q	L	E	A	M	I	R	F	L	T	R	K	D	P	D	L	E	C	D	E	
L	V	R	E	V	L	F	D	A	V	V	T	A	P	M	E	A	Y	W	T	A	L	A	L	N	M	S	E	E	
E	H	V	V	D	M	A	F	L	G	T	R	A	G	L	L	R	S	S	L	F	V	G	S	E	K	V	S	D	
K	F	L	T	P	E	D	E	A	S	V	F	T	L	D	R	F	P	L	W	Y	R	Q	A	S	E	H	P	A	
S	F	V	F	N	L	R	W	A	E	G	P	E	S	A	G	E	P	M	V	V	T	A	S	T	A	V	A		
V	D	K	R	T	A	I	A	A	A	G	V	Q	M	K	L	E	F	L	Q	R	K	F	W	A	A	T	R	Q	
C	S	T	V	D	G	P	Y	T	Q	S	C	E	D	S	D	L	D	C	F	V	I	D	N	N	G	F	I	L	
S	K	R	S	R	E	T	G	R	F	L	G	E	V	D	G	A	V	L	T	Q	L	L	S	M	G	V	F	S	
V	T	M	Y	D	Y	Q	A	M	C	K	P	S	S	H	H	H	S	A	A	Q	P	L	V	S	P	I	S	A	
L	T	A	T	R	W	L	L	Q	E	L	V	L	F	L	L	E	W	S	V	W	G	S	W	Y	D	R	G		
A	K	S	V	F	H	H	S	H	K	H	K	K	Q	D	P	L	Q	P	C	D	T	E	Y	P	V	F	V	Y	
P	A	I	R	E	A	N	G	I	V	E	C	G	P	C	Q	K	V	F	V	Q	Q	I	P	N	S	N	L	K	
L	L	V	T	D	P	T	C	D	C	S	I	F	P	P	V	L	Q	E	A	T	E	V	K	Y	N	A	S	V	
C	D	R	M	R	S	Q	K	L	R	R	R	P	D	S	C	H	A	F	H	P	E	V	R	V	E	A	D	R	
W	A	G	F	S	S	P	N	P	L	C	L	G	L	C	P	C	R	Q	E	H	I	G	M	P	M	N	T	P	
P	V	L	L	G	G	N	I	R	V	Y	A	L																	

Comparison of ha2δ-4.pep with ha2δ3.pep:

April 7, 2000 14:19

Percent Similarity: 69.172 Percent Identity: 60.133

1MAVALGTRRRDR.....VKLWADTFG	21		
:				
1	MAGPGSPRRASRGASALLAALLYAALGDVVRSEQQIPLSVV	KLWASAFG	50	
22	GDLYNTVTKYSGSLLLQKKYKDVESSLKIEEV	DGLELVRKFSED	MENMLR	71
: .			: . : : : . .	
51	GEIKSIAAKYSGSQLLQKKYKEYKDVAIEIDGLQLV	KKLA	NMEEMFH	100

846 SISCDDETVCYLIIDNNNGFILVSEDYTQTGdffGEIEGAVMNKLLTMGSF 895
 .
 866 SQVTMYDYQAMCKPSSHHSAAQPLVSPISAFLTATRWLLQELVLFLEW 915
 .
 896 KRITLYDYQAMCRANKESSDGAHGLDPYNAFLSAVKWIMTELVLFLVEF 945
 .
 916 SVWGSWYDRGAEAKSVFHHSHKHKKQDPLQPCDTEYPVFVYQPAIREANG 965
 .
 946 NLC.SWWHSMDTAKA.....QKLKQTLLEPCDTEYPAFVSERTIKETTG 987
 .
 966 IVECGPCQKVFFVQQIPNSNLLLVTDPDCSIFPPVQLQEATEVKYNAS 1015
 .
 988 NIACEDCSKSFVIQQIPSSNLFMVVVDSSCLCESVAPITMAPIEIRYNES 1037
 .
 1016 VKCDRMRSQKLRRRPDSCHAFHPEVRVEADRGWAGFSSPN.....PLCLG 1060
 .
 1038 LKCERLKAQKIRRRPESCHGFHPEENARECGGAPSLQAQTVLLLLPLLLM 1087
 .
 1061 LCPCRQEHI GMPMNTPVPVLLGGNIRVYAL 1090
 |
 1088 LFSR..... 1091

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